

IN THE CLAIMS

Please amend the claims as follows:

Claim 1 (Original): A chimeric protein derived from a high-threshold calcium channel, characterized in that it comprises at least one  $\beta$  subunit or a fragment thereof including at least the BID domain, fused, at its NH<sub>2</sub> or COON end, with the I-II loop of an  $\alpha_1$  subunit or fragment thereof including at least the AID domain.

Claim 2 (Original): The chimeric protein as claimed in claim 1, characterized in that it consists of a  $\beta$  subunit fused, at its NH<sub>2</sub> or at its COOH end, with the I-II loop of an  $\alpha_1$  subunit.

Claim 3 (Original): The chimeric protein as claimed in claim 1, characterized in that it consists of the GK-like domain of a  $\beta$  subunit fused, at its NH<sub>2</sub> or COOH end, with the I-II loop of an  $\alpha_1$  subunit.

Claim 4 (Currently Amended): The protein of claim 1 ~~as claimed in any one of claims 1 to 3~~, characterized in that the  $\beta$  subunit, or a fragment thereof, and the I-II loop, or a fragment thereof, are separated by a spacer peptide.

Claim 5 (Currently Amended): The chimeric protein of claim 1 ~~as claimed in any one of claims 1 to 4~~, characterized in that it is derived from a G-protein-sensitive high-threshold calcium channel.

Claim 6 (Original): The chimeric protein as claimed in claim 5, characterized in that it comprises the I-II loop of an  $\alpha_1$  subunit selected from  $\alpha_{1A}$ ,  $\alpha_{1B}$  and  $\alpha_{1E}$ , or a fragment thereof.

Claim 7 (Currently Amended): The chimeric protein as claimed of claim 1 ~~in any one of claims 1 to 6~~, characterized in that it comprises a  $\beta$  subunit selected from the group consisting of  $\beta_1$ ,  $\beta_2$ ,  $\beta_3$  and  $\beta_4$ , or a fragment thereof.

Claim 8 (Currently Amended): A variant chimeric protein derived from a chimeric protein as claimed in claim 1 ~~any one of claims 1 to 7~~, characterized in that it has a mutation of at least one amino acid in the sequences of said  $\beta$  subunit and/or of the I-II loop of an  $\alpha_1$  subunit.

Claim 9 (Original): The variant chimeric protein as claimed in claim 8, characterized in that said mutation modifies the affinity of the  $\beta$  subunit for the fragment of the I-II loop of the  $\alpha$  subunit and/or vice versa.

Claim 10 (Currently Amended): The variant chimeric protein as claimed in claim 8 ~~or claim 9~~, characterized in that said mutations are selected from the following mutations of the AID domain of the I-II loop of the  $\alpha_1$  subunit: Q383A, Q384A, E386D, E386S, L389H, G391R, Y392S, Y392F, W395A, I396A and E400A.

Claim 11 (Currently Amended): The chimeric protein as claimed in ~~any one of claims 1 to 10~~ claim 1, characterized in that it is coupled, ~~preferably covalently~~, to at least one suitable label allowing the detection and/or the purification and/or the immobilization of said protein.

Claim 12 (Original): The chimeric protein as claimed in claim 11, characterized in that it comprises an acceptor or donor fluorophore respectively at its NH<sub>2</sub> and/or COOH end.

Claim 13 (Original): The chimeric protein as claimed in claim 12, characterized in that the acceptor fluorophore is the fluorescent protein CFP or BFP and the donor fluorophore is the fluorescent protein GFP or YFP.

Claim 14 (Currently Amended): A peptide, characterized in that it comprises a fragment of at least 7 amino acids of the sequence of the chimeric protein as claimed in claim 1 ~~any one of claims 1 to 13~~, which fragment includes at least the 7 amino acids located at the junction of the  $\beta$  subunit and of the I-II loop of the  $\alpha_1$  subunit of a calcium channel ~~or of their fragments as defined in claim 1~~.

Claim 15 (Currently Amended): An antibody, characterized in that it is directed against a peptide as claimed in claim 14, ~~and in that it recognizes exclusively the chimeric protein as claimed in any one of claims 1 to 13~~.

Claim 16 (Currently Amended): A nucleic acid molecule, characterized in that it is selected from ~~the group consisting of the sequences encoding a chimeric protein as claimed in any one of claims 1 to 13 or a peptide as claimed in claim 14~~ claim 1, and the sequences complementary to the above sequences, that may be sense or antisense.

Claim 17 (Original): Probes and primers, characterized in that they comprise a sequence of approximately 10 to 30 nucleotides corresponding to that located at the junction of the  $\beta$  subunit and of the I-II loop of the  $\alpha_1$  subunit of a calcium channel or of their fragments as defined in claim 1.

Claim 18 (Original): Primers capable of amplifying the  $\beta$  subunit and/or the I-II loop of the  $\alpha_1$  subunit of a calcium channel or their fragments as defined in claim 1, characterized in that they are selected from the group consisting of the sequences SEQ ID NO: 1, 2, 4, 6, 7, 8 and 9.

Claim 19 (Original): A recombinant vector, characterized in that it comprises an insert selected from the group consisting of the nucleic acid molecules as claimed in claim 16.

Claim 20 (Original): The recombinant vector as claimed in claim 19, characterized in that it is a eukaryotic expression vector having a sequence selected from the group consisting of the sequences SEQ ID NO: 5 and SEQ ID NO: 10.

Claim 21 (Currently Amended): A cell modified with a recombinant vector as claimed in claim 19 or 20, ~~with a nucleic acid molecules as claimed in claim 16 or with a chimeric protein as claimed in any one of claims 1 to 13.~~

Claim 22 (Original): The modified cell as claimed in claim 21, characterized in that it is a eukaryotic cell.

Claim 23 (Currently Amended): The modified cell as claimed in claim 21 ~~or claim 22~~, characterized in that it expresses at least one receptor capable of coupling to G proteins.

Claim 24 (Original): A nonhuman transgenic mammal, characterized in that all or some of its cells are transformed with a nucleic acid molecule as claimed in claim 16.

Claim 25 ( Currently Amended ): The use of a the product ~~selected from the group consisting of the chimeric proteins as claimed in any one of claims 1 to 13, the nucleic acid molecules as claimed in claim 16, the recombinant vectors as claimed in claim 19 or claim 20, the modified cells as claimed in any one of claims 21 to 23 and the nonhuman transgenic mammals as claimed in claim 24~~ of claim 1[[,]] for studying the G-protein-coupled receptor-dependent cell signaling and regulatory pathways.

Claim 26 (Currently Amended): The use of a the product selected from ~~the group consisting of the chimeric proteins as claimed in any one of claims 1 to 13, the nucleic acid molecules as claimed in claim 16, the recombinant vectors as claimed in claim 19 or claim 20, the modified cells as claimed in any one of claims 21 to 23 and the nonhuman transgenic mammals as claimed in claim 24~~ claim 1[[,]] for screening agonists and/or antagonists of G-protein-coupled receptor-dependent cell signaling and regulatory pathways.

Claim 27 (Currently Amended): The use of a the product selected from ~~the group consisting of the chimeric proteins as claimed in any one of claims 1 to 13, the nucleic acid molecules as claimed in claim 16, the recombinant vectors as claimed in claim 19 or claim 20, the modified cells as claimed in any one of claims 21 to 23 and the nonhuman transgenic mammals as claimed in claim 24~~ of claim 1, for screening antagonists of the interaction between the  $\alpha_1$  and  $\beta$  subunits of high-threshold calcium channels.

Claim 28 (Original): A method for studying the G-protein-coupled receptor-dependent cell signaling and regulatory pathways, characterized in that it comprises at least the following steps:

- a) culturing of modified cells expressing a chimeric protein derived from a G-protein-sensitive calcium channel and a G-protein-coupled receptor, as claimed in claim 23,
- b) transduction of a signal via said G-protein-coupled receptor, by any appropriate means, and
- c) determination, by any appropriate means, of the proportion of said chimeric protein expressed in said cells that is bound to a  $G\beta\gamma$  subunit.

Claim 29 (Original): A method for screening agonists/antagonists of the G-protein-coupled receptor-dependent cell signaling and regulatory pathways, characterized in that it comprises at least the following steps:

- a<sub>2</sub>) culturing of modified cells expressing a chimeric protein derived from a G-protein-sensitive calcium channel and a G-protein-coupled receptor, as claimed in claim 23,
- b<sub>2</sub>) transduction of a signal via said G-protein-coupled receptor, by any appropriate means,
- c<sub>2</sub>) comparative determination, by any appropriate means, of the proportion of said chimeric protein expressed in the cells that is bound to a Gβγ subunit, before and after the bringing into contact of said cells in b<sub>2</sub>) with a molecule to be tested, and
- d<sub>2</sub>) identification of the molecules that are agonists/antagonists of the G-protein-coupled receptor-dependent cell signaling and regulatory pathways, corresponding to those capable respectively of increasing and of decreasing the cellular concentration of free Gβγ subunits.

Claim 30 (Currently Amended): The method as claimed in claim 28 ~~or claim 29~~, characterized in that said modified cells in a<sub>1</sub>) or in a<sub>2</sub>) express a chimeric protein coupled, at its NH<sub>2</sub> and COOH ends, respectively to a fluorescence donor fluorophore and a fluorescence acceptor fluorophore, and said determination in c<sub>1</sub>) or in c<sub>2</sub>) is carried out by means of the fluorescence transfer (FRET) technique.

Claim 31 (Currently Amended): A method for screening antagonists of the interaction between the  $\alpha_1$  and  $\beta$  subunits of high-threshold calcium channels, characterized in that it comprises at least the following steps:

- a<sub>3</sub>) bringing a molecule to be tested into contact with a chimeric protein derived from a G-protein-sensitive or -insensitive calcium channel as claimed ~~in any one of claims 1 to 13~~ claim 1 and with a peptide comprising the AID domain of a G-protein-insensitive  $\alpha$  subunit,
- b<sub>3</sub>) measuring, by any appropriate means, the binding of said chimeric protein to said peptide, and
- c<sub>3</sub>) identifying the antagonists of the interaction between the  $\alpha_1$  and  $\beta$  subunits corresponding to those with which binding of said chimeric protein to said peptide is observed.

Claim 32 (Currently Amended): The screening method as claimed in claim 31, characterized in that said peptide comprising the AID domain is immobilized on a solid support and said chimeric protein is a chimeric protein ~~as claimed in any one of claims 11 to 13.~~

Claim 33 (Currently Amended): A kit for implementing a method as claimed in any ~~one of claims 28 to 32~~ claim 28, characterized in that it comprises at least one product selected from the group consisting of the chimeric proteins ~~as claimed in any one of claims 1 to 13~~, the nucleic acid molecules ~~as claimed in claim 16~~, the recombinant vectors ~~as claimed in claim 19 or claim 20~~, the modified cells ~~as claimed in any one of claims 21 to 23~~ and the nonhuman transgenic mammals ~~as claimed in claim 24~~.